<u>RAW SEQUENCE LISTING</u>

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/559,952_
Source: _/FwP_
Date Processed by STIC: _12/20/05_

ENTERED

IFWP

RAW SEQUENCE LISTING   DATE: 12/20/2005
PATENT APPLICATION: US/10/559,952   TIME: 10:25:16

Input Set : A:\1003 SEQLIST.txt
Output Set: N:\CRF4\12202005\J559952.raw

```
   4 <110> APPLICANT: Roche, Andrew
   5         Hansen, Martin Chr.
   6         Villsen, Inge D.
   7         Schrotz-King, Petra
   8         Henningsen, Jeannette
   9         Lund Jorgensen, Trine Louise
  11 <120> TITLE OF INVENTION: Extracellular Aspergillus Polypeptides
  14 <130> FILE REFERENCE: 13403.1003
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/559,952
C--> 16 <141> CURRENT FILING DATE: 2005-12-09
  16 <160> NUMBER OF SEQ ID NOS: 49
  18 <170> SOFTWARE: FastSEQ for Windows Version 4.0
  20 <210> SEQ ID NO: 1
  21 <211> LENGTH: 260
  22 <212> TYPE: PRT
  23 <213> ORGANISM: Aspergillus Fumigatus
  25 <400> SEQUENCE: 1
  26 Met Leu Ala Ser Phe Gln Phe Cys Ile Leu Pro Arg Thr Tyr Arg Thr
  27 1               5                   10                  15
  28 Leu Leu Cys Ser Ala Gly Ala Gly Pro Leu Leu Ile Ile Gln Phe Val
  29             20                  25                  30
  30 Thr Val Ala Ser Ala Leu Ala Leu Ala Pro Thr Ala Val Val Ala Arg
  31             35                  40                  45
  32 Gln Gly Ala Ala Ala Phe Val Thr Val Asn Ser Ile Asp Val Cys Pro
  33         50                  55                  60
  34 Lys Lys Val Ala Gln Glu Ile Ile Asn Pro Gly Pro Lys Val Val Thr
  35 65                  70                  75                  80
  36 Thr Pro Tyr Thr Cys Asp Gln Val Lys Leu Gly His Gly Leu Asp Val
  37             85                  90                  95
  38 Ser Tyr Tyr Asn Phe Asp Ile Glu Pro Leu Thr Lys Asp Thr Phe Pro
  39             100                 105                 110
  40 Tyr Cys Lys Ala Leu Lys Val Phe Asp Asn Glu Gly Cys Leu Gly Phe
  41         115                 120                 125
  42 Pro Thr Leu Trp Ile Pro Leu Glu Ser Pro Leu Glu Asp Lys Cys Ile
  43     130                 135                 140
  44 Pro Glu His Tyr Phe Ser Asp Glu Val Lys Ser Ile Ser Phe Gln Leu
  45 145                 150                 155                 160
  46 Asp Cys Arg Glu Asp Ala Pro Val Lys Lys Glu Pro Tyr Gly Pro Lys
  47             165                 170                 175
  48 Glu Gly Ala Glu Gln Ser Ala Pro Gln Ala Glu His Ser Thr Lys Gln
  49             180                 185                 190
  50 Asp Ala Gln Gln Gly Ser His Gln Gly Gln Glu Val Gln Asn Ser Pro
  51             195                 200                 205
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52 Lys Gln Glu Ala Arg Gln Gly Ser Arg Pro Ala Glu Ala Ala Pro Lys
53     210               215               220
54 Gln Glu Gln Glu Ala Glu Gln Ala Ser Glu Ala Ala Pro Glu Lys Lys
55 225               230               235               240
56 Ala Ser Asn Pro Ala Asp Ser Leu Gly Leu Gly Glu Leu Thr Lys Val
58               245               250               255
59 Leu Gly Phe Arg
60               260
62 <210> SEQ ID NO: 2
63 <211> LENGTH: 107
64 <212> TYPE: PRT
65 <213> ORGANISM: Aspergillus Fumigatus
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68 Val Arg Phe Pro Val Pro Asp Asp Ile Thr Val Lys Gln Ala Thr Glu
69 1               5               10              15
70 Lys Cys Gly Asp Gln Ala Gln Leu Ser Cys Cys Asn Lys Ala Thr Tyr
71             20              25              30
72 Ala Gly Asp Val Thr Asp Ile Asp Glu Gly Ile Leu Ala Gly Thr Leu
73             35              40              45
74 Lys Asn Leu Ile Gly Gly Gly Ser Gly Thr Glu Gly Leu Gly Leu Phe
75     50              55              60
76 Asn Gln Cys Ser Lys Leu Asp Leu Gln Ser Pro Ile Ile Gly Ile Pro
77 65              70              75              80
78 Ile Gln Asp Leu Val Asn Gln Lys Cys Lys Gln Asn Ile Ala Cys Cys
79             85              90              95
80 Gln Asn Ser Pro Ser Asp Ala Val Arg Phe Pro
81             100             105
83 <210> SEQ ID NO: 3
84 <211> LENGTH: 318
85 <212> TYPE: PRT
86 <213> ORGANISM: Aspergillus Fumigatus
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91 Ile Val Gly Leu Asn Ser Leu Ser His Gly Val Asp Val Val Ala Val
92             20              25              30
93 Asn Asp Pro Phe Ile Glu Val His Tyr Ala Ala Tyr Met Leu Lys Tyr
94             35              40              45
95 Asp Thr Thr His Gly Gln Phe Lys Gly Thr Ile Glu Thr Tyr Asp Gln
96     50              55              60
97 Gly Leu Ile Val Asn Gly Lys Lys Ile Arg Phe Tyr Ala Glu Lys Asp
98 65              70              75              80
99 Pro Ser Gln Ile Pro Trp Ser Glu Thr Gly Ala Ala Tyr Ile Val Glu
100             85              90              95
101 Ser Thr Gly Val Phe Thr Thr Lys Glu Lys Ala Ser Ala His Leu Lys
102             100             105             110
103 Gly Gly Ala Lys Lys Val Ile Ile Ser Ala Pro Ser Ala Asp Ala Pro
104             115             120             125
105 Met Phe Val Met Gly Val Asn Asn Thr Thr Tyr Thr Ser Asp Ile Gln
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106      130                    135                    140
107 Val Leu Ser Asn Ala Ser Cys Thr Thr Asn Cys Leu Ala Pro Leu Ala
108 145                    150                    155                    160
109 Lys Val Ile Asn Asp Lys Phe Gly Ile Val Glu Gly Leu Met Thr Thr
110                    165                    170                    175
111 Val His Ser Tyr Thr Ala Thr Gln Lys Val Val Asp Ala Pro Ser Asn
112                    180                    185                    190
113 Lys Asp Trp Arg Gly Gly Arg Thr Ala Ala Gln Asn Ile Ile Pro Ser
114             195                    200                    205
115 Ser Thr Gly Ala Ala Lys Ala Val Gly Lys Val Ile Pro Ser Leu Asn
116        210                    215                    220
117 Gly Lys Leu Thr Gly Met Ala Met Arg Val Pro Thr Ser Asn Val Ser
118 225                    230                    235                    240
119 Val Val Asp Leu Thr Cys Arg Leu Glu Lys Gly Ala Ser Tyr Asp Glu
120                    245                    250                    255
121 Ile Lys Gln Ala Ile Lys Ala Ala Ser Glu Glu Gly Glu Leu Lys Asn
122                    260                    265                    270
123 Ile Leu Gly Tyr Thr Glu Asp Asp Val Val Ser Ser Asp Leu Asn Gly
124             275                    280                    285
125 Asp Glu Arg Ser Ser Ile Phe Asp Ala Lys Ala Gly Ile Ser Leu Asn
126        290                    295                    300
127 Pro Asn Phe Val Lys Leu Val Ala Trp Tyr Asp Asn Glu Trp
128 305                    310                    315
130 <210> SEQ ID NO: 4
131 <211> LENGTH: 438
132 <212> TYPE: PRT
133 <213> ORGANISM: Aspergillus Fumigatus
135 <400> SEQUENCE: 4
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138 Asn Pro Thr Val Glu Val Asp Val Ala Thr Glu Thr Gly Leu His Arg
139                    20                    25                    30
140 Ala Ile Val Pro Ser Gly Ala Ser Thr Gly Gln His Glu Ala His Glu
141        35                    40                    45
142 Leu Arg Asp Gly Asp Lys Thr Gln Trp Gly Gly Lys Gly Val Leu Lys
143        50                    55                    60
144 Ala Val Lys Asn Val Asn Glu Thr Ile Gly Pro Ala Leu Ile Lys Glu
145 65                    70                    75                    80
146 Asn Ile Asp Val Lys Asp Gln Ser Lys Val Asp Glu Phe Leu Asn Lys
147                    85                    90                    95
148 Leu Asp Gly Thr Ala Asn Lys Ser Asn Leu Gly Ala Asn Ala Ile Leu
149                    100                    105                    110
150 Gly Val Ser Leu Ala Val Ala Lys Ala Gly Ala Ala Glu Lys Gly Val
151        115                    120                    125
152 Pro Leu Tyr Ala His Ile Ser Asp Leu Ala Gly Thr Lys Lys Pro Tyr
153        130                    135                    140
154 Val Leu Pro Val Pro Phe Gln Asn Val Leu Asn Gly Gly Ser His Ala
155 145                    150                    155                    160
156 Gly Gly Arg Leu Ala Phe Gln Glu Phe Met Ile Val Pro Asp Ser Ala
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157                    165                  170                  175
158 Pro Ser Phe Ser Glu Ala Leu Arg Gln Gly Ala Glu Val Tyr Gln Lys
159                    180                  185                  190
160 Leu Lys Ala Leu Ala Lys Lys Lys Tyr Gly Gln Ser Ala Gly Asn Val
161         195                  200                  205
162 Gly Asp Glu Gly Gly Val Ala Pro Asp Ile Gln Thr Ala Glu Glu Ala
163     210                  215                  220
164 Leu Asp Leu Ile Thr Glu Ala Ile Glu Gln Ala Gly Tyr Thr Gly Lys
165 225                  230                  235                  240
166 Ile Lys Ile Ala Met Asp Val Ala Ser Ser Glu Phe Tyr Lys Ala Asp
167                    245                  250                  255
168 Val Lys Lys Tyr Asp Leu Asp Phe Lys Asn Pro Glu Ser Asp Pro Ser
169         260                  265                  270
170 Lys Trp Leu Thr Tyr Glu Gln Leu Ala Asp Leu Tyr Lys Ser Leu Ala
171         275                  280                  285
172 Ala Lys Tyr Pro Ile Val Ser Ile Glu Asp Pro Phe Ala Glu Asp Asp
173     290                  295                  300
174 Trp Glu Ala Trp Ser Tyr Phe Tyr Lys Thr Ser Asp Phe Gln Ile Val
175 305                  310                  315                  320
176 Gly Asp Asp Leu Thr Val Thr Asn Pro Gly Arg Ile Lys Lys Ala Ile
177                    325                  330                  335
178 Glu Leu Lys Ser Cys Asn Ala Leu Leu Leu Lys Val Asn Gln Ile Gly
179         340                  345                  350
180 Thr Leu Thr Glu Ser Ile Gln Ala Ala Lys Asp Ser Tyr Ala Asp Asn
181         355                  360                  365
182 Trp Gly Val Met Val Ser His Arg Ser Gly Glu Thr Glu Asp Val Thr
183     370                  375                  380
184 Ile Ala Asp Ile Ala Val Gly Leu Arg Ser Gly Gln Ile Lys Thr Gly
185 385                  390                  395                  400
186 Ala Pro Cys Arg Ser Glu Arg Leu Ala Lys Leu Asn Gln Ile Leu Arg
187             405                  410                  415
188 Ile Glu Glu Glu Leu Gly Glu Asn Thr Val Tyr Ala Gly Ser Lys Phe
189         420                  425                  430
190 Arg Thr Ala Val Asn Leu
191         435
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 728
195 <212> TYPE: PRT
196 <213> ORGANISM: Aspergillus Fumigatus
198 <400> SEQUENCE: 5
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201 Cys Pro Tyr Met Thr Gly Glu Leu Asn Arg Arg Asp Glu Ile Ser Asp
202         20                  25                  30
203 Gly Asp Ala Ala Ala Ala Thr Glu Glu Phe Leu Ser Gln Tyr Tyr Leu
204     35                  40                  45
205 Asn Asp Asn Asp Ala Phe Met Thr Ser Asp Val Gly Gly Pro Ile Glu
206     50                  55                  60
207 Asp Gln Asn Ser Leu Ser Ala Gly Glu Arg Gly Pro Thr Leu Leu Glu
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208 65                      70                  75                      80
209 Asp Phe Ile Phe Arg Gln Lys Ile Gln Arg Phe Asp His Glu Arg Val
210                     85                  90                      95
211 Pro Glu Arg Ala Val His Ala Arg Gly Ala Gly Ala His Gly Val Phe
212                     100                 105                     110
213 Thr Ser Tyr Gly Asp Phe Ser Asn Ile Thr Ala Ala Ser Phe Leu Ala
214         115                 120                     125
215 Lys Glu Gly Lys Gln Thr Pro Val Phe Val Arg Phe Ser Thr Val Ala
216     130                 135                     140
217 Gly Ser Arg Gly Ser Ser Asp Leu Ala Arg Asp Val His Gly Phe Ala
218 145                 150                     155                     160
219 Thr Arg Phe Tyr Thr Asp Glu Gly Asn Phe Asp Ile Val Gly Asn Asn
220                 165                 170                     175
221 Ile Pro Val Phe Phe Ile Gln Asp Ala Ile Leu Phe Pro Asp Leu Ile
222         .       180                 185                     190
223 His Ala Val Lys Pro Arg Gly Asp Asn Glu Ile Pro Gln Ala Ala Thr
224         . 195                 200                     205
225 Ala His Asp Ser Ala Trp Asp Phe Phe Ser Gln Gln Pro Ser Thr Met
226     210                 215                     220
227 His Thr Leu Leu Trp Ala Met Ser Gly His Gly Ile Pro Arg Ser Phe
228 225                 230                     235                     240
229 Arg His Val Asp Gly Phe Gly Val His Thr Phe Arg Phe Val Thr Asp
230                 245                 250                     255
231 Asp Gly Ala Ser Lys Leu Val Lys Phe His Trp Lys Ser Leu Gln Gly
232                 260                 265                     270
233 Lys Ala Ser Met Val Trp Glu Glu Ala Gln Gln Thr Ser Gly Lys Asn
234         275                 280                     285
235 Pro Asp Phe Met Arg Gln Asp Leu His Asp Ala Ile Glu Ala Gly Arg
236     290                 295                     300
237 Tyr Pro Glu Trp Glu Leu Gly Val Gln Ile Met Asp Glu Glu Asp Gln
238 305                 310                 315                     320
239 Leu Arg Phe Gly Phe Asp Leu Leu Asp Pro Thr Lys Ile Val Pro Glu
240                 325                 330                     335
241 Glu Phe Val Pro Ile Thr Lys Leu Gly Lys Met Gln Leu Asn Arg Asn
242                 340                 345                     350
243 Pro Arg Asn Tyr Phe Ala Glu Thr Glu Gln Val Met Phe Gln Pro Gly
244         355                 360                     365
245 His Ile Val Arg Gly Val Asp Phe Thr Glu Asp Pro Leu Leu Gln Gly
246     370                 375                     380
247 Arg Leu Phe Ser Tyr Leu Asp Thr Gln Leu Asn Arg His Gly Gly Pro
248 385                 390                     395                     400
249 Asn Phe Glu Gln Leu Pro Ile Asn Gln Pro Arg Val Pro Val His Asn
250                 405                 410                     415
251 Asn Asn Arg Asp Gly Ala Gly Gln Met Phe Ile Pro Leu Asn Pro His
252         420                 425                     430
253 Ala Tyr Ser Pro Lys Thr Ser Val Asn Gly Ser Pro Lys Gln Ala Asn
254         435                 440                     445
255 Gln Thr Val Gly Asp Gly Phe Phe Thr Ala Pro Gly Arg Thr Thr Ser
256     450                 455                     460
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/20/2005
PATENT APPLICATION: **US/10/559,952**   TIME: 10:25:17

Input Set : **A:\1003 SEQLIST.txt**
Output Set: **N:\CRF4\12202005\J559952.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the
<220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:36; Xaa Pos. 176,179
Seq#:37; Xaa Pos. 2,5
Seq#:38; N Pos. 579,580,581,588,589,590
Seq#:39; N Pos. 526,527,528,535,536,537

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/559,952

DATE: 12/20/2005
TIME: 10:25:17

Input Set : A:\1003 SEQLIST.txt
Output Set: N:\CRF4\12202005\J559952.raw

L:16 M:270 C: Current Application Number differs, Replaced Current Application No
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:160
L:724 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:176
L:760 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:540
L:815 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:480